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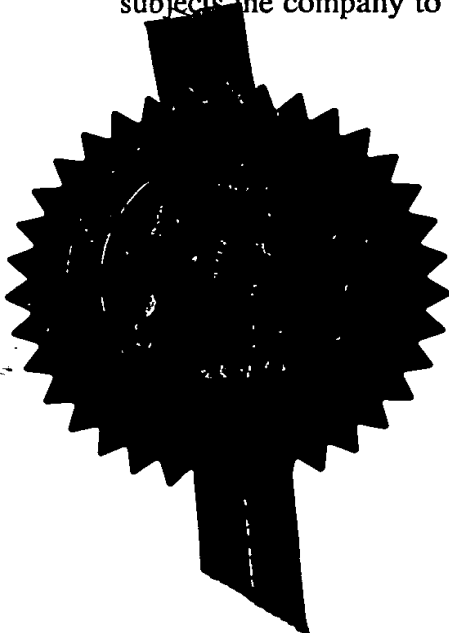
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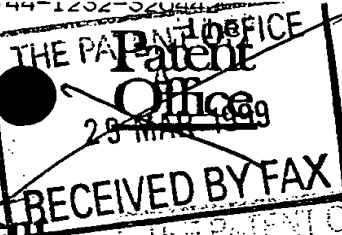
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Abbreviations: Area under the curve, AUC; Dipeptidyl peptidase IV, DPP IV; Electrospray ionization mass spectrometry, ESI-MS; Gastric inhibitory polypeptide, GIP; glucagon-like peptide-1(7-36)amide, tGLP-1; Trifluoroacetic acid, TFA.

Gastric inhibitory polypeptide (GIP) is an important insulin-releasing hormone of the enteroinsular axis which like glucagon-like peptide-1(7-36)amide (tGLP-1) has a functional profile of possible therapeutic value for NIDDM. Both incretin hormones are rapidly inactivated in plasma by the exopeptidase dipeptidyl peptidase IV (DPP IV). The present study has examined the ability of N-terminal modification of human GIP to protect from plasma degradation and enhance insulin-releasing and antihyperglycemic activity. Degradation of GIP by incubation at 37°C with purified DPP IV was clearly evident after 4 h (54% intact). After 12 h, more than 60% of GIP was converted to GIP(3-42) whereas >99% N-terminally modified Tyr¹-glucitol GIP remained intact. Tyr¹-glucitol GIP was similarly resistant to serum degradation. The formation of GIP(3-42) was almost completely abolished by inhibition of plasma DPP IV with diprotin A. Effects of GIP and Tyr¹-glucitol GIP were examined in Wistar rats following i.p. injection of either peptide (10 nmol/kg) together with glucose (18 mmol/kg). Plasma glucose concentrations were significantly lower and insulin concentrations higher following both peptides compared with glucose alone. More importantly, individual glucose values at 15 min and 30 min together with the areas under the curve (AUC) for glucose were significantly lower following administration of Tyr¹-glucitol GIP as compared to GIP (AUC, 255±33 versus 368±8 mmol/l.min, respectively; $P<0.01$). This was associated with a significantly greater and more protracted insulin response following Tyr¹-glucitol GIP than GIP (AUC, 773±41 versus 639±39 ng/ml.min; $P<0.05$). These data demonstrate that Tyr¹-glucitol GIP displays resistance to plasma DPP IV degradation and exhibits enhanced antihyperglycemic activity and insulin-releasing action in vivo.



INTRODUCTION

Gastric inhibitory polypeptide (GIP) and glucagon-like peptide-1(7-36)amide (truncated GLP-1; tGLP-1) are two important insulin-releasing hormones secreted from endocrine cells in the intestinal tract in response to feeding [1,2]. Together with autonomic nerves they play a vital supporting role to the pancreatic islets in the control of blood glucose homeostasis and nutrient metabolism [1,3].

Dipeptidyl peptidase IV (DPP IV; EC 3.4.14.5) has been identified as a key enzyme responsible for inactivation of GIP and tGLP-1 in serum [4,5]. DPP IV is completely inhibited in serum by the addition of diprotin A (DPA, 0.1 mmol/l) [4]. This occurs through the rapid removal of the N-terminal dipeptides Tyr¹-Ala² and His⁷-Ala⁸ giving rise to the main metabolites GIP(3-42) and GLP-1(9-36)amide, respectively. These truncated peptides are reported to lack biological activity or to even serve as antagonists at GIP or tGLP-1 receptors [6-9]. The resulting biological half-lives of these incretin hormones in vivo are therefore very short, estimated to be no longer than approximately 5 min, respectively [5,10-12]. In situations of normal glucose regulation and pancreatic B-cell sensitivity, this short duration of action is advantageous in facilitating momentary adjustments to homeostatic control. However, the current goal of a possible therapeutic role of incretin hormones, particularly tGLP-1 in NIDDM therapy is frustrated by a number of factors in addition to finding a convenient route of administration [13]. Most notable of these are rapid peptide degradation and rapid absorption (peak concentrations reached 20 min) and the resulting need for both high dosage and precise timing with meals [13-15]. Recent therapeutic strategies have focussed on precipitated preparations to delay peptide absorption [16] and inhibition of GLP-1 degradation using specific inhibitors of

DPP IV [17-19]. A possible therapeutic role is also suggested by the observation that a specific inhibitor of DPP IV, isoleucine thiazolidide, lowered blood glucose and enhanced insulin secretion in glucose-treated diabetic obese Zucker rats presumably by protecting against catabolism of the incretin hormones tGLP-1 and GIP [18].

Numerous studies have indicated that tGLP-1 infusion restores pancreatic B-cell sensitivity, insulin secretory oscillations and improved glycemic control in various groups of patients with IGT or NIDDM [13,15,20-22]. Longer term studies also show significant benefits of tGLP-1 injections in NIDDM and possibly IDDM therapy [20,23,24], providing a major incentive to develop an orally effective or long-acting tGLP-1 analogue [13]. Several attempts have been made to produce structurally modified analogues of tGLP-1 which are resistant to DPP IV degradation [25-27]. A significant extension of serum half-life is observed with His⁷-glucitol tGLP-1 and tGLP-1 analogues substituted at position 8 with Gly, Aib, Ser or Thr [25-27]. However, these structural modifications appear to impair receptor binding and insulinotropic activity thereby compromising the part of the benefits of protection from proteolytic degradation [25-28]. Thus in our own recent studies using His⁷-glucitol tGLP-1, resistance to DPP IV and serum degradation was accompanied by severe loss of insulin-releasing activity [26,28].

GIP shares not only the same degradation pathway as tGLP-1 but many similar physiological actions, including stimulation of insulin and somatostatin secretion, enhancement of glucose disposal [1]. These actions are viewed as key aspects in the antihyperglycemic properties of tGLP-1 [13], and there is therefore good expectation that GIP may have similar potential in NIDDM therapy. Indeed, compensation by GIP is held to explain the modest disturbances of

glucose homeostasis observed in tGLP-1 knockout mice [29]. Apart from early studies [30], the anti-diabetic potential of GIP has not been explored and tGLP-1 may seem more attractive since it is viewed by some as a more potent insulin secretagogue when infused at 'so called' physiological concentrations estimated by RIA [31].

In a recent study, we have shown that N-terminal glycation of GIP markedly enhances the insulin releasing effect of the peptide on clonal B-cells [32]. If such structural modification also confers DPP IV resistance, the potential attractiveness of this peptide for possible NIDDM therapy would be considerable enhanced. The present study has explored this issue by examining in vitro degradation of Tyr¹-glucitol GIP together with evaluation of its antihyperglycemic and insulin-releasing properties in vivo. The results demonstrate clearly that this novel GIP analogue exhibits a substantial resistance to aminopeptidase degradation and increased glucose lowering activity compared with the native human GIP.

RESEARCH DESIGN AND METHODS

Materials. Human GIP was purchased from the American Peptide Company (Sunnyvale, CA, USA). HPLC grade acetonitrile was obtained from Rathburn (Walkersburn, Scotland). Sequencing grade trifluoroacetic acid (TFA) was obtained from Aldrich (Poole, Dorset, UK). All other chemicals purchased including dextran T-70, activated charcoal, sodium cyanoborohydride and bovine serum albumin fraction V were from Sigma (Poole, Dorset, UK). Diprotin A (DPA) was purchased from Calbiochem-Novabiochem (UK) Ltd. (Beeston, Nottingham, UK) and rat insulin standard for RIA was obtained from Novo

Industria (Copenhagen, Denmark). Reversed-phase Sep-Pak cartridges (C-18) were purchased from Millipore-Waters (Milford, MA, USA). All water used in these experiments was purified using a Milli-Q Water Purification System (Millipore Corporation, Milford, MA, U.S.A.).

Preparation of Tyr¹-glucitol GIP. Tyr¹-glucitol GIP was prepared and purified by HPLC as described previously [32]. In brief, human GIP was incubated with glucose under reducing conditions in 10 mmol/l sodium phosphate buffer at pH 7.4 for 24 h. The reaction was stopped by addition of 0.5 mol/l acetic acid (30 µl) and the mixture applied to a Vydac (C-18) (4.6 x 250 mm) analytical HPLC column (The Separations Group, Hesperia, CA, USA) and gradient elution conditions were established using aqueous/TFA and acetonitrile/TFA solvents, as described previously. Fractions corresponding to the glycosylated peaks were pooled, taken to dryness under vacuum using an AES 1000 Speed-Vac concentrator (Life Sciences International, Runcorn, UK) and purified to homogeneity on a Supelcosil (C-8) (4.6 x 150 mm) column (Supelco Inc., Poole Dorset, UK).

Degradation of GIP and Tyr¹-glucitol GIP by DPP IV. HPLC-purified GIP or Tyr¹-glucitol GIP were incubated at 37°C with DPP-IV (5 mU) for various time periods in a reaction mixture made up to 500 µl with 50 mmol/l triethanolamine-HCl, pH 7.8 (final peptide concentration 1 µmol/l) [4]. Enzymatic reactions were terminated after 0, 2, 4 and 12 h by addition of 5 µl of 10% (v/v) TFA/water. Samples were made up to a final volume of 1.0 ml with 0.12% (v/v) TFA and stored at -20°C prior to HPLC analysis.

Degradation of GIP and Tyr¹-glucit 1 GIP by human plasma. Pooled human plasma (20 μ l) taken from six healthy fasted male subjects was incubated at 37°C with GIP or Tyr¹-glucitol GIP (10 μ g) for 0 and 4 h in a reaction mixture made up to 500 μ l, containing 50 mmol/l triethanolamine/HCl buffer pH 7.8. Incubations for 4 h were also performed in the presence of diprotin A (5 mU). The reactions were terminated by addition of 5 μ l of TFA and the final volume adjusted to 1.0 ml using 0.1% v/v TFA/water. Samples were centrifuged (13,000g, 5 min) and the supernatant applied to a C-18 Sep-Pak cartridge (Millipore-Waters) which was previously primed and washed with 0.1% (v/v) TFA/water. After washing with 20 ml 0.12% TFA/water, bound material was released by elution with 2 ml of 80% (v/v) acetonitrile/water and concentrated using a Speed-Vac concentrator (Runcorn, UK). The volume was adjusted to 1.0 ml with 0.12% (v/v) TFA/water prior to HPLC purification.

HPLC analysis of degraded GIP and Tyr¹-glucitol GIP. Samples were applied to a Vydac C-18 widepore column equilibrated with 0.12% (v/v) TFA/H₂O at a flow rate of 1.0 ml/min. Using 0.1% (v/v) TFA in 70% acetonitrile/H₂O, the concentration of acetonitrile in the eluting solvent was raised from 0% to 31.5% over 15 min, to 38.5% over 30 min and from 38.5% to 70% over 5 min, using linear gradients. The absorbance was monitored at 206 nm and peak areas valuated using a model 2221 LKB integrator. Samples recovered manually were concentrated using a Speed-Vac concentrator.

Electrospray ionization mass spectrometry (ESI-MS). Samples for ESI-MS analysis containing intact and degradation fragments of GIP (from DPP IV and

plasma incubations) as well as Tyr¹-glucitol GIP, were further purified by HPLC. Peptides were dissolved (approximately 400 pmol) in 100 µl of water and applied to the LCQ benchtop mass spectrometer (Finnigan MAT, Hemel Hempstead, UK) equipped with a microbore C-18 HPLC column (150 x 2.0 mm, Phenomenex, UK, Ltd., Macclesfield). Samples (30 µl direct loop injection) were injected at a flow rate of 0.2 ml/min, under isocratic conditions 35% (v/v) acetonitrile/water. Mass spectra were obtained from the quadrupole ion trap mass analyzer and recorded. Spectra were collected using full ion scan mode over the mass-to-charge (m/z) range 150-2000. The molecular masses of GIP and related structures were determined from ESI-MS profiles using prominent multiple charged ions and the following equation $M_r = iM_i - iM_h$ (where M_r = molecular mass; M_i = m/z ratio; i = number of charges; M_h = mass of a proton).

In vivo biological activity of GIP and Tyr¹-glucitol GIP. Effects of GIP and Tyr¹-glucitol GIP on plasma glucose and insulin concentrations were examined using 10-12 week old male Wistar rats. The animals were housed individually in an air conditioned room at 22±2°C with a 12 h light/12 h dark cycle. Drinking water and a standard rodent maintenance diet (Trouw Nutrition, Belfast) were supplied ad libitum. Food was withdrawn for an 18 h period prior to intraperitoneal injection of glucose alone (18 mmol/kg body weight) or in combination with either GIP or Tyr¹-glucitol GIP (10 nmol/kg). Test solutions were administered in a final volume of 8 ml/kg body weight. Blood samples were collected at 0, 15, 30 and 60 min from the cut tip of the tail of conscious rats into chilled fluoride/heparin microcentrifuge tubes (Sarstedt, Nümbrecht, Germany). Samples were centrifuged using a Beckman microcentrifuge for 30

sec at 13,000 g. Plasma samples were aliquoted and stored at -20°C prior to glucose and insulin determinations. All animal studies were done in accordance with the Animals (Scientific Procedures) Act 1986.

Analyses. Plasma glucose was assayed by an automated glucose oxidase procedure using a Beckman Glucose Analyzer II [33]. Plasma insulin was determined by dextran charcoal radioimmunoassay as described previously [34]. Incremental areas under plasma glucose and insulin curves (AUC) were calculated using a computer program (CAREA) employing the trapezoidal rule [35] with baseline subtraction. Results are expressed as mean \pm SEM and values were compared using the Student's unpaired *t*-test. Groups of data were considered to be significantly different if $P < 0.05$.

RESULTS

Degradation of GIP and Tyr¹-glucitol GIP by DPP IV. Fig. 1 illustrates the typical peak profiles obtained from the HPLC separation of the products obtained from the incubation of GIP (left panels) or Tyr¹-glucitol GIP (right panels) with DPP IV for 0, 2, 4 and 12 h. The retention times of GIP and Tyr¹-glucitol GIP at $t=0$ were 21.93 min and 21.75 min, respectively. Degradation of GIP was evident after 4 h incubation (54% intact), and by 12 h the majority (60%) of intact GIP was converted to the single product with a retention time of 21.61 min. Tyr¹-glucitol GIP remained almost completely intact throughout 2-12 h incubation.

Degradation of GIP and Tyr¹-glucitol GIP by human plasma. Fig. 2 shows a set of typical HPLC profiles of the products obtained from the incubation of GIP or Tyr¹-

glucitol GIP with human plasma for 0 and 4 h. GIP (left panels) with a retention time of 22.06 min was readily metabolised by plasma within 4 h incubation giving rise to the appearance of a major degradation peak with a retention time of 21.74 min. In contrast, the incubation of Tyr¹-glucitol GIP under similar conditions (right panels) did not result in the formation of any detectable degradation fragments during this time with only a single peak being observed with a retention time of 21.77 min. Addition of diprotin A, a specific inhibitor of DPP IV, to GIP during the 4 h incubation completely inhibited degradation of the peptide by plasma.

Identification of GIP degradation fragments by ESI-MS. Fig. 3 shows the monoisotopic molecular masses obtained for GIP, (panel A), Tyr¹-glucitol GIP (panel B) and the major plasma degradation fragment of GIP (panel C) using ESI-MS. The peptides analyzed were purified from plasma incubations as shown in Fig. 2. The exact molecular mass (M_r) of the peptides were calculated using the equation $M_r = iM_l - iM_h$ as defined in Research Design and Methods section. After spectral averaging was performed, prominent multiple charged species $(M+3H)^{3+}$ and $(M+4H)^{4+}$ were detected from GIP at m/z 1661.6 and 1246.8, corresponding to intact M_r 4981.8 and 4983.2 Da, respectively (Fig. 3A). Similarly, for Tyr¹-glucitol GIP $((M+4H)^{4+}$ and $(M+5H)^{5+})$ were detected at m/z 1287.7 and 1030.3, corresponding to intact molecular masses of M_r 5146.8 and 5146.5 Da, respectively (Fig. 3B). The difference between the observed molecular masses of the quadruply charged GIP and the N-terminally modified GIP species (163.6 Da) indicated that the latter peptide contained a single glucitol adduct corresponding to Tyr¹-glucitol GIP. Fig. 3C shows the prominent multiply charged species

($M+3H$)³⁺ and ($M+4H$)⁴⁺ detected from the major fragment of GIP at m/z 1583.8 and 1188.1, corresponding to intact M_r 4748.4 and 4748.4 Da, respectively. This corresponds with the theoretical mass of the N-terminally truncated form of the peptide GIP(3-42). This fragment was also the major degradation product of DPP IV incubations (data not shown).

Effects of GIP and Tyr¹-glucitol GIP on plasma glucose homeostasis. Fig. 4-5 show the effects of i.p. glucose alone (18 mmol/kg) (control group), and glucose in combination with GIP or Tyr¹-glucitol GIP (10 nmol/kg) on plasma glucose and insulin concentrations. Compared with the control group, plasma glucose concentrations and area under the curve (AUC) were significantly lower following administration of either GIP or Tyr¹-glucitol GIP (Fig. 4A, B). Furthermore, individual values at 15 and 30 min together with AUC were significantly lower following administration of Tyr¹-glucitol GIP as compared to GIP. Consistent with the established insulin-releasing properties of GIP, plasma insulin concentrations of both peptide-treated groups were significantly raised at 15 and 30 min compared with the values after administration of glucose alone (Fig. 5A). The overall insulin responses, estimated as AUC were also significantly greater for the two peptide-treated groups (Fig. 5B). Despite lower prevailing glucose concentrations than the GIP-treated group, plasma insulin response, calculated as AUC, following Tyr¹-glucitol GIP was significantly greater than after GIP (Fig. 5B). The significant elevation of plasma insulin at 30 min is of particular note, suggesting that the insulin-releasing action of Tyr¹-glucitol GIP is more protracted than GIP even in the face of a diminished glycemic stimulus (Fig. 4A, 5A).

DISCUSSION

The forty-two amino acid GIP is an important incretin hormone released into the circulation from endocrine K-cells of the duodenum and jejunum following ingestion of food [36]. The high degree of structural conservation of GIP among species supports the view that this peptide plays an important role in metabolism [12]. Secretion of GIP is stimulated directly by actively transported nutrients in the gut lumen without a notable input from autonomic nerves [12]. The most important stimulants of GIP release are simple sugars [37] and unsaturated long chain fatty acids [38], with amino acids exerting weaker effects [39]. As with tGLP-1, the insulin-releasing effect of GIP is strictly glucose-dependent [30,40]. This affords protection against hypoglycemia and thereby fulfils one of the most desirable features of any current or potentially new antidiabetic drug [41].

The present results demonstrate for the first time that Tyr¹-glucitol GIP displays profound resistance to serum and DPP IV degradation. Using ESI-MS the present study showed that native GIP was rapidly cleaved in vitro to a major 4748.4 Da degradation product, corresponding to GIP(3-42) which confirmed previous findings using matrix-assisted laser desorption ionization time-of-flight mass spectrometry [42]. Serum degradation was completely inhibited by diprotin A (Ile-Pro-Ile), a specific competitive inhibitor of DPP IV, confirming this as the main enzyme for GIP inactivation in vivo [4,5]. In contrast, Tyr¹-glucitol GIP remained almost completely intact after incubation with serum or DPP IV for up to 12 h. This indicates that glycation of GIP at the amino-terminal Tyr¹ residue masks the potential cleavage site from DPP IV and prevents removal of the Tyr¹-Ala² dipeptide from the N-terminus preventing the formation of GIP(3-42).

Consistent with in vitro protection against DPP IV, administration of Tyr¹-

glucitol GIP significantly enhanced the antihyperglycemic activity and insulin-releasing action of the peptide when administered with glucose to rats. Native GIP enhanced insulin release and reduced the glycemic excursion as observed in many previous studies [12,40]. However, amino-terminal glycation of GIP increased the insulin-releasing and antihyperglycemic actions of the peptide by 62% and 38% respectively, as estimated from AUC measurements. Detailed kinetic analysis is difficult due to necessary limitation of sampling times, but the greater insulin concentrations following Tyr¹-glucitol GIP as opposed to GIP at 30 min post-injection is indicative of longer half-life. The glycemic rise was modest in both peptide-treated groups and glucose concentrations following injection of Tyr¹-glucitol GIP were consistently lower than after GIP. Since the insulinotropic actions of GIP are glucose-dependent [30,40], it is likely that the relative insulin-releasing potency of Tyr¹-glucitol GIP is greatly underestimated in the present in vivo experiments.

In keeping with this interpretation, recent in vitro studies in our laboratory using glucose-responsive clonal B-cells showed that the insulin-releasing potency of Tyr¹-glucitol GIP was several orders of magnitude greater than GIP and that its effectiveness was more sensitive to change of glucose concentrations within the physiological range [32]. Together with the present in vivo observations, this suggests that N-terminal glycation of GIP confers resistance to DPP IV degradation while enhancing receptor binding and insulin secretory effects on the B-cell. These attributes of Tyr¹-glucitol GIP are fully expressed in vivo where DPP IV resistance impedes degradation of the peptide to GIP(3-42), thereby prolonging the half-life and enhancing effective concentrations of the intact biologically active peptide. It is thus possible that glycated GIP is

enhancing insulin secretion in vivo both by enhanced potency at the receptor as well as improving DPP IV resistance. Thus numerous studies have shown that GIP(3-42) and other N-terminally modified fragments, including GIP(4-42), and GIP(17-42) are either weakly effective or inactive in stimulating insulin release [4,43-45]. Furthermore, evidence exists that N-terminal deletions of GIP result in receptor antagonist properties in GIP receptor transfected Chinese hamster kidney cells [9], suggesting that inhibition of GIP catabolism would also reduce the possible feedback antagonism at the receptor level by the truncated GIP(3-42).

In addition to its insulintropic actions, a number of other potentially important extrapancreatic actions of GIP may contribute to the enhanced antihyperglycemic activity and other beneficial metabolic effects of Tyr¹-glucitol GIP. These include the stimulation of glucose uptake in adipocytes, increased synthesis of fatty acids and activation of lipoprotein lipase in adipose tissue [46-48]. GIP also promotes plasma triglyceride clearance in response to oral fat loading [49]. In liver, GIP has been shown to enhance insulin-dependent inhibition of glycogenolysis [50]. GIP also reduces both glucagon-stimulated lipolysis in adipose tissue as well as hepatic glucose production [51]. Finally, recent findings indicate that GIP has a potent effect on glucose uptake and metabolism in mouse isolated diaphragm muscle [52]. This latter action may be shared with tGLP-1 [53,54] and both peptides have additional benefits of stimulating somatostatin secretion and slowing down gastric emptying and nutrient absorption [1,55].

In conclusion, this study has demonstrated for the first time that the glycation of GIP at the amino-terminal Tyr¹ residue limits GIP catabolism through impairment of the proteolytic actions of serum peptidases and thus

prolongs its half-life in vivo. This effect is accompanied by enhanced antihyperglycemic activity and raised insulin concentrations in vivo, suggesting that such DPP IV resistant analogues should be explored alongside tGLP-1 as potentially useful therapeutic agents for NIDDM. Tyr¹-glucitol GIP appears to be particularly interesting in this regard since such amino-terminal modification of GIP enhances [32] rather than impairs glucose-dependent insulintropic potency as was observed recently for tGLP-1 [28].

Acknowledgements

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Legends t Figures

Fig. 1. Degradation of GIP and Tyr¹-glucitol GIP by DPP IV. Representative HPLC profiles obtained after incubation of GIP (left panels) or Tyr¹-glucitol GIP (right panels) with DPP IV for 0, 2, 4 and 12 h. Incubations of GIP and Tyr¹-glucitol GIP exposed to DPP IV were separated on a Vydac C-18 column using linear gradients 0% to 31.5% acetonitrile over 15 min, to 38.5% over 30 min and from 38.5% to 70% acetonitrile over 5 min. Left hand panels show HPLC profiles of intact GIP (retention time 21.93 min) and GIP(3-42) (retention time 21.61 min). Right hand panels show HPLC profiles obtained for Tyr¹-glucitol GIP (retention time 21.75 min). HPLC peaks corresponding to intact GIP, GIP(3-42) and Tyr¹-glucitol GIP are indicated.

Fig. 2. Degradation of GIP and Tyr¹-glucitol GIP by human plasma. Representative HPLC profiles obtained after incubation of GIP (left panels) and Tyr¹-glucitol GIP (right panels) with human plasma for 0 and 4 h and for 4 h in the presence of 5 mU of diprotin A (DPA). GIP and Tyr¹-glucitol GIP incubations were separated with a Vydac C-18 column using linear gradients 0% to 31.5% acetonitrile over 15 min, to 38.5% over 30 min and from 38.5% to 70% acetonitrile over 5 min. Peaks corresponding with intact GIP, GIP(3-42) and Tyr¹-glucitol GIP are indicated. A major peak corresponding to the specific DPP IV inhibitor tripeptide DPA appears in the bottom panels with retention time 16.29 min.

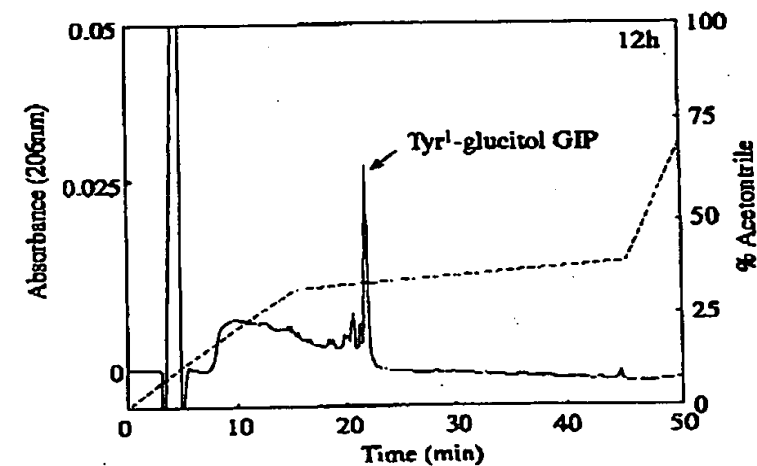
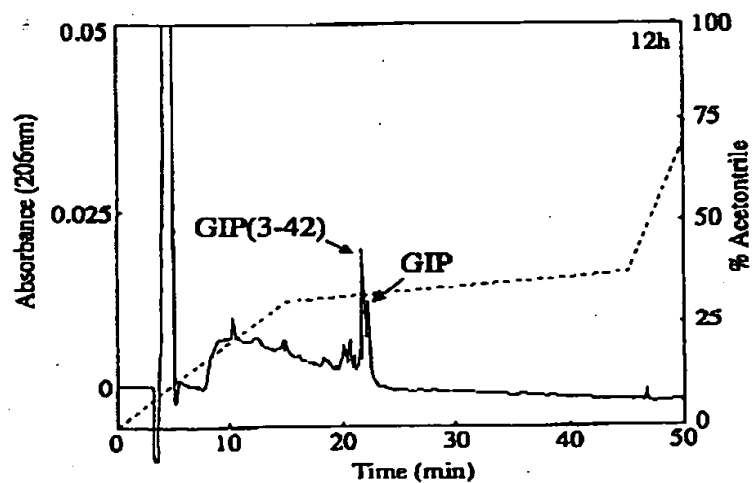
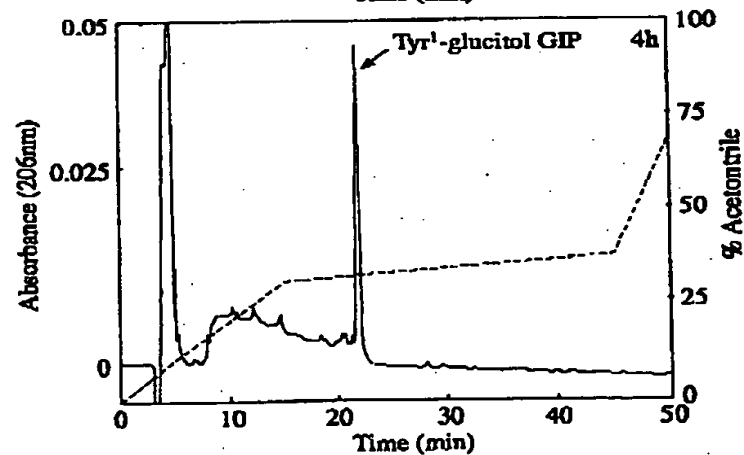
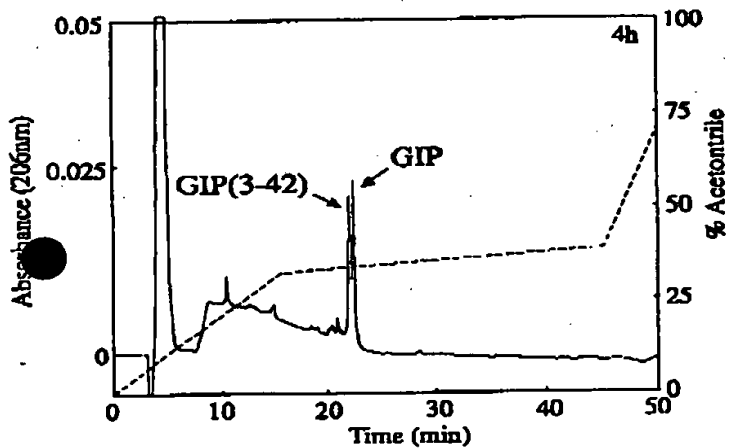
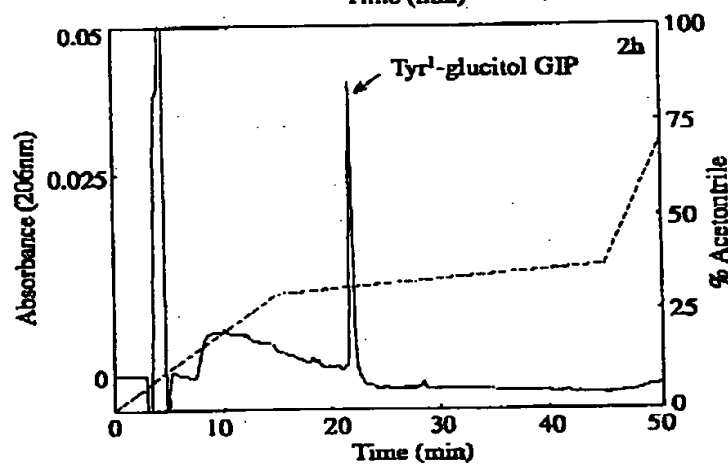
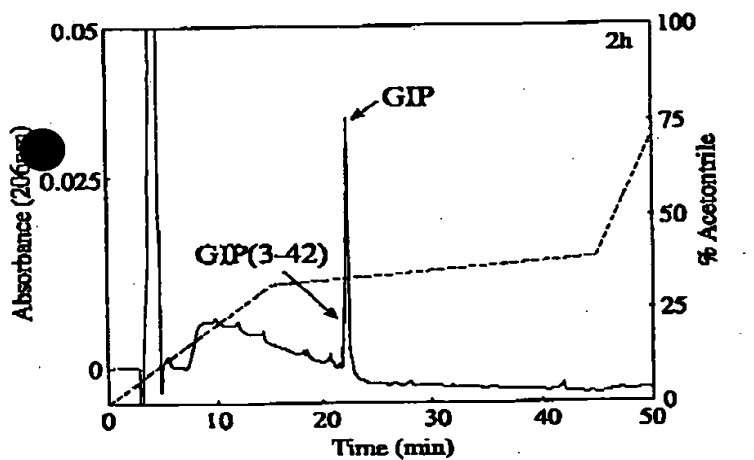
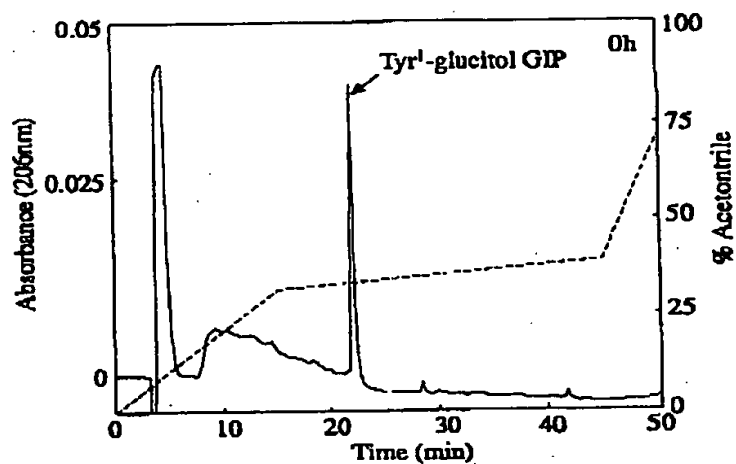
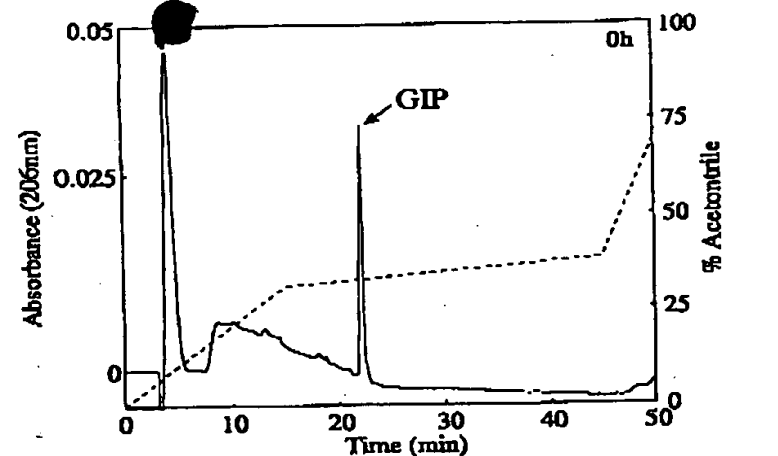
Fig. 3. Electrospray ionization mass spectrometry of GIP, Tyr1-glucitol GIP and the major degradation fragment GIP(3-42). Samples containing (A) GIP, (B) Tyr1-glucitol GIP and (C) the major degradation fragment of GIP (GIP(3-42)) isolated from plasma incubations (Fig. 2). Peptides were dissolved (approximately 400 pmol) in 100µl of water and applied to the LC/MS equipped with a microbore C-18 HPLC column. Samples (30 µl direct loop injection) were applied at a flow rate of 0.2 ml/min, under isocratic conditions 35% acetonitrile/water. Mass spectra were recorded using a quadripole ion trap mass analyzer. Spectra were collected using full ion scan mode over the mass-to-charge (m/z) range 150-2000. The molecular masses (M_r) of GIP and related structures were determined from ESI-MS profiles using prominent multiple charged ions and the following equation $M_r = iM_i - iM_h$ (see Research Design and Methods section).

Fig. 4. Effects of GIP and glycated GIP on plasma glucose homeostasis. (A) Plasma glucose concentrations after i.p. glucose alone (18 mmol/kg) (control group), or glucose in combination with either GIP or Tyr1-glucitol GIP (10 nmol/kg). The time of injection is indicated by the arrow (0 min). (B) Plasma glucose AUC values for 0-60 min post injection. Values are mean \pm SEM for six rats. ** $P < 0.01$, *** $P < 0.001$ compared with GIP and Tyr1-glucitol GIP; + $P < 0.05$, ++ $P < 0.01$ compared with non-glycated GIP.

Fig. 5. Effects of GIP on plasma insulin responses. (A) Plasma insulin concentrations after i.p. glucose alone (18 mmol/kg) (control group), or glucose in combination with either with GIP or glycated GIP (10 nmol/kg). The time of injection is indicated by the arrow. **(B)** Plasma insulin AUC values were calculated for each of the 3 groups up to 90 min post injection. The time of injection is indicated by the arrow (0 min). Plasma insulin AUC values for 0-60 min post injection. Values are mean \pm SEM for six rats. * $P < 0.05$, ** $P < 0.01$ *** $P < 0.001$ compared with GIP and Tyr¹-glucitol GIP; + $P < 0.05$, ++ $P < 0.01$ compared with non-glycated GIP.

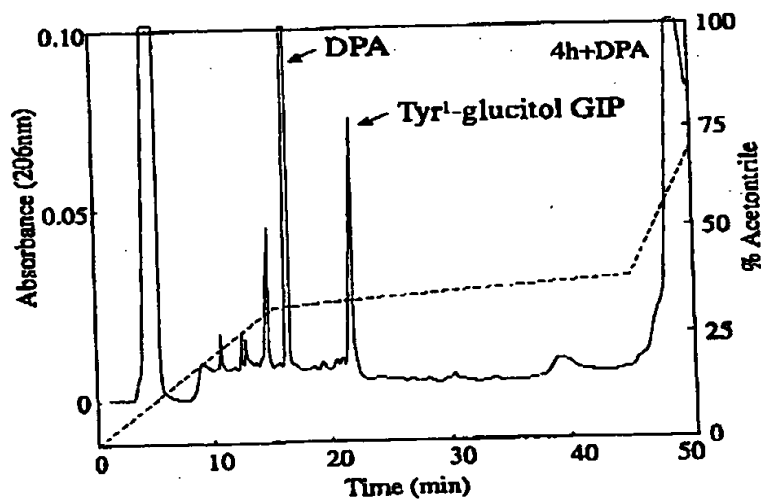
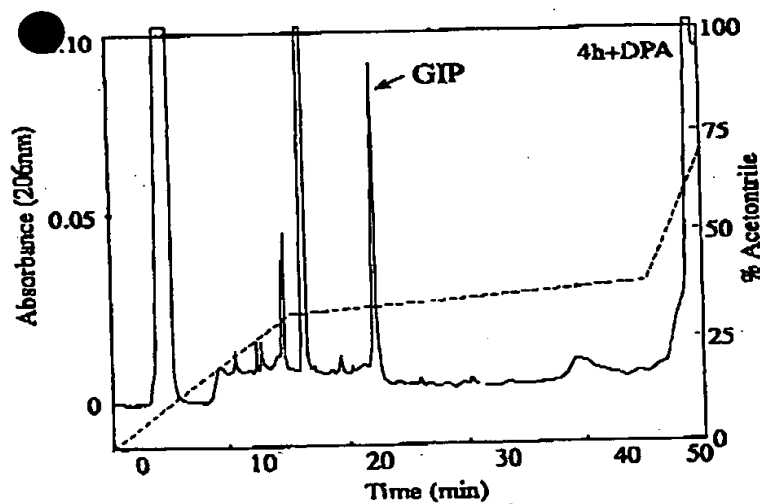
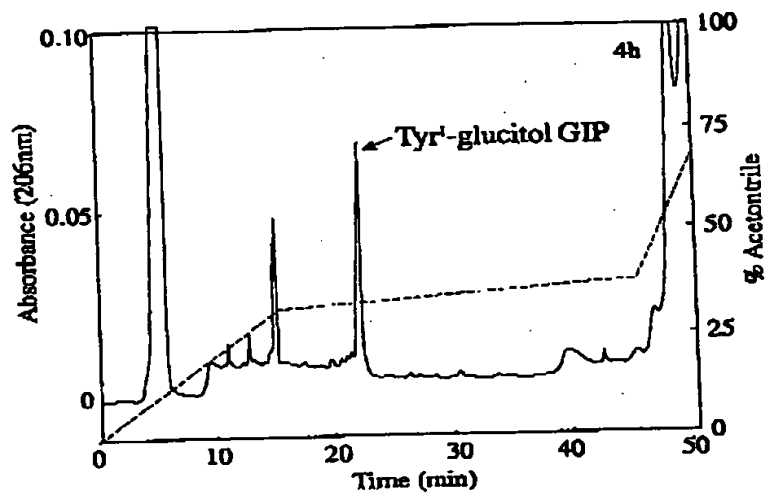
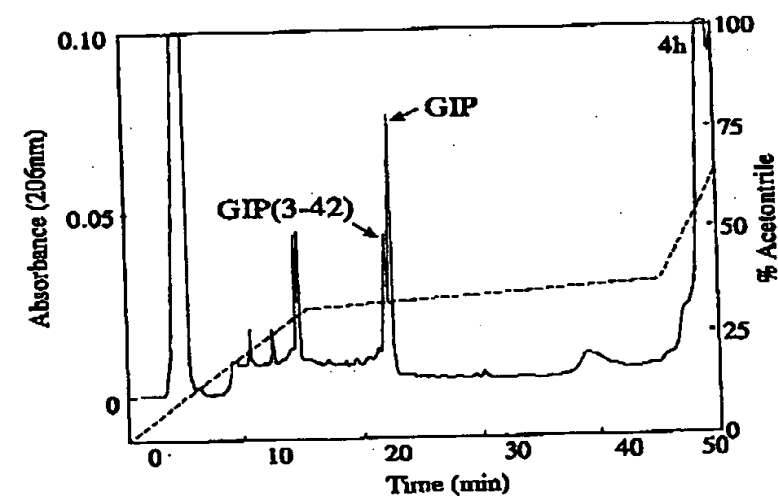
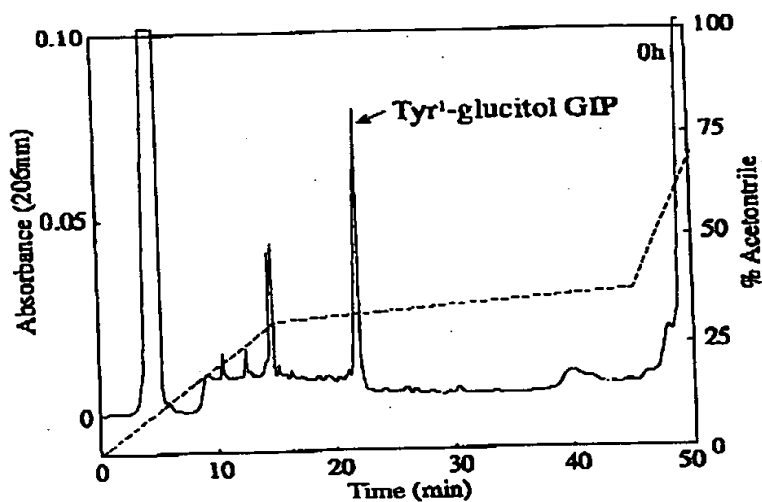
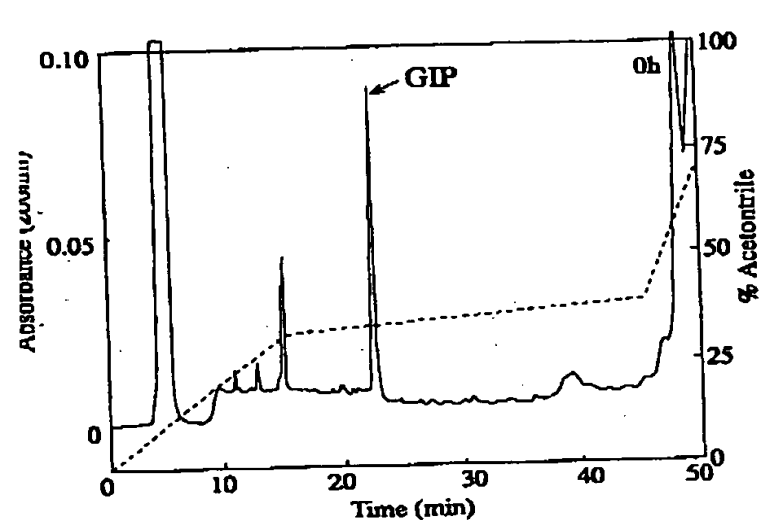
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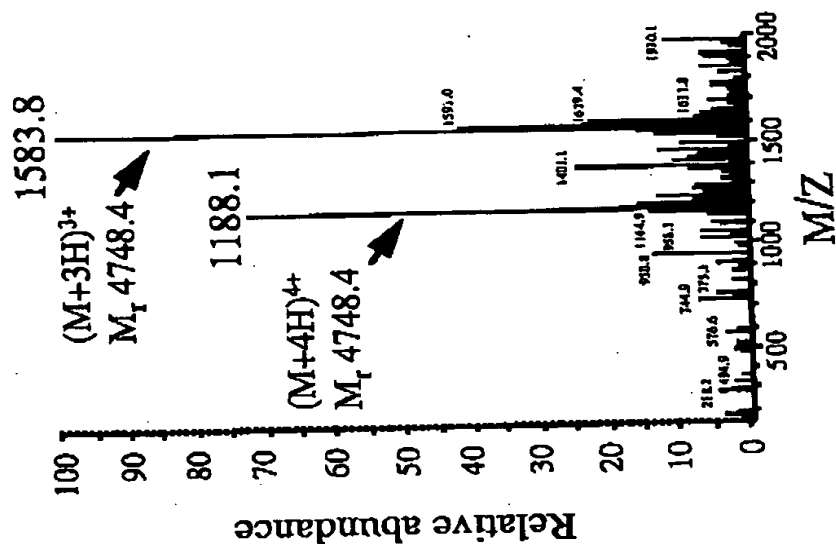
Fig 2



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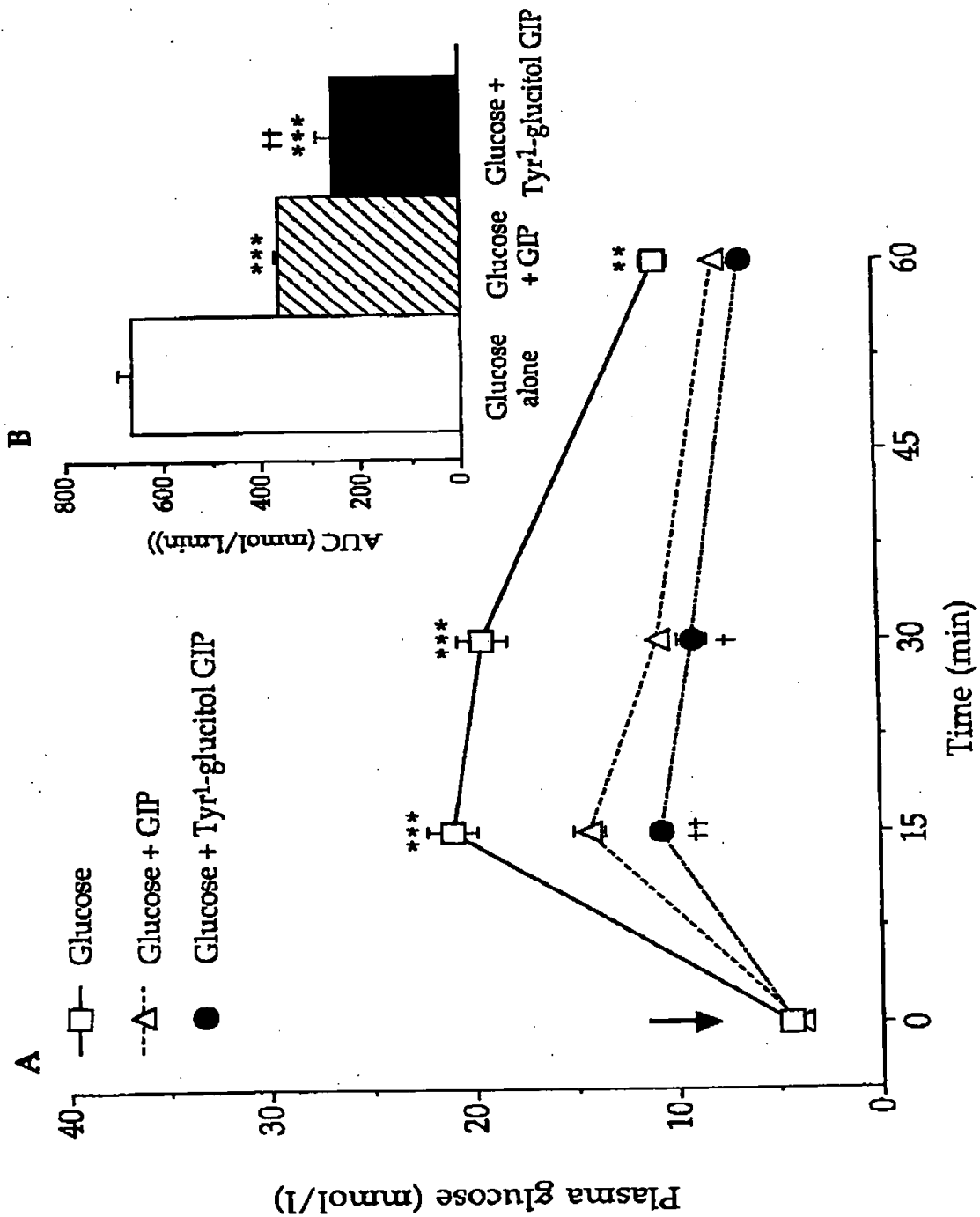
Fig 3

(C) GIP(3-42)



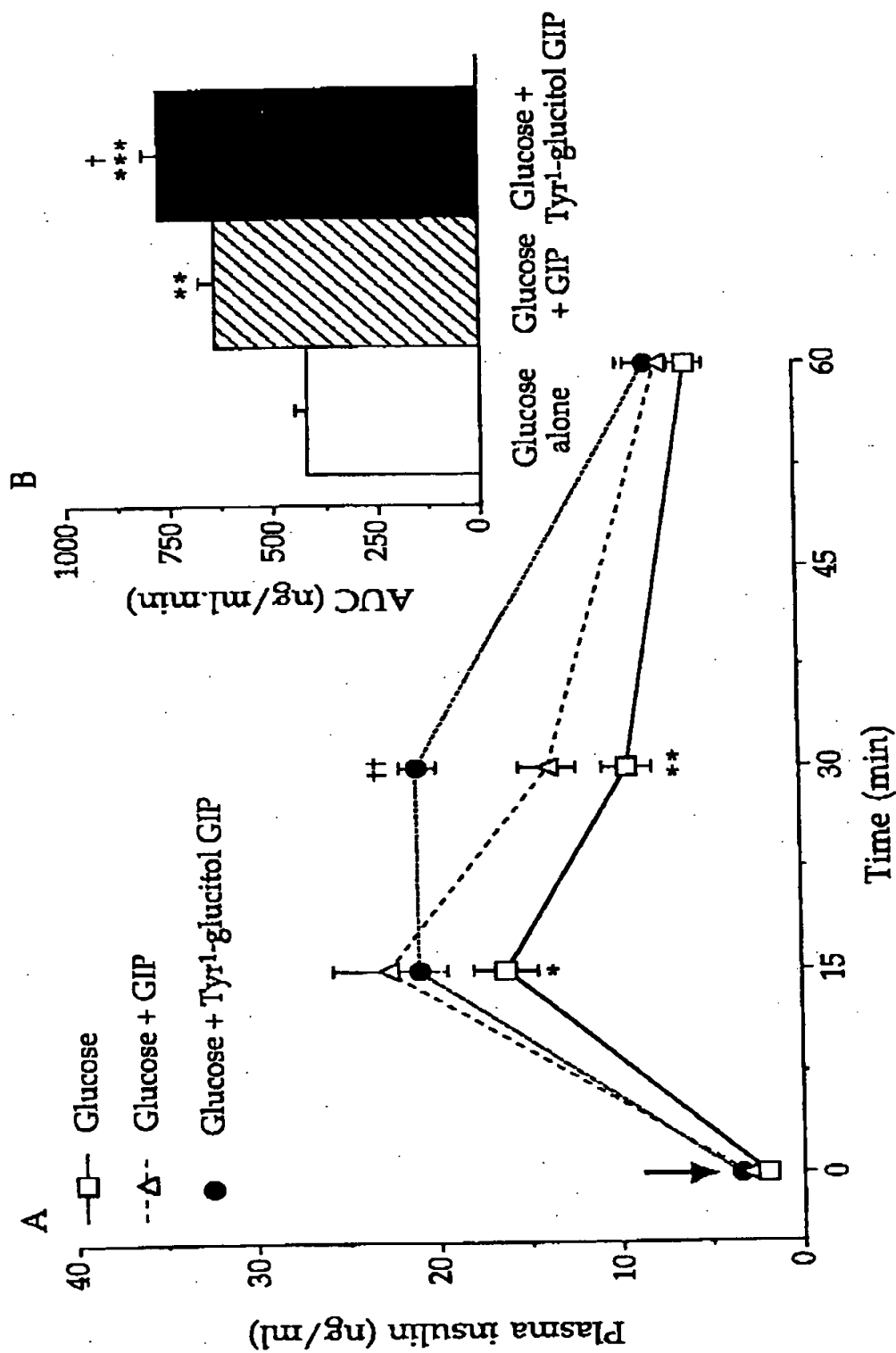
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Fig 4



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Fig 5



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